Amendments to the Claims:

This listing of claims will replace all prior versions, and listings of claims in the application:

Listing of Claims:

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1 1. (Currently Amended): A method for determining the presence of a microbial 2 organism an organism of interest in a sample from another organism or organisms to be 3 distinguished; said method comprising: 4 treating the sample, or a portion thereof, with at least one detectable molecular 5 probe wherein the molecular probe or probes are peptide nucleic acid and are selected 6 such that either: 7 a target sequence of both the microbial organism of interest and the other (i) 8 organism or organisms reacts with the molecular probe in a way that 9 produces detectable microbial organisms of interest and a detectable other 10 organism or organisms to be distinguished; or 11 a target sequence of only the microbial organism of interest reacts with the (ii) molecular probe in a way that produces only detectable organisms of 12 13 interest: and 14 contacting the sample, or a portion thereof, with a solid carrier to which has been 15 immobilized an antibody binding partner such that if (i) applies then the antibody binding 16 partner is chosen to be reactive only with the detectable microbial organism of interest 17 but not reactive with the detectable other organism or organisms to be distinguished; but 18 if (ii) applies then the <u>antibody</u> binding partner is chosen to be generally reactive with the 19 detectable microbial organism of interest but also may be reactive with the other 20 organism or organisms to be distinguished; and 21 determining the presence, absence, position or number of detectable microbial 22 organisms immobilized to the solid carrier and correlating the result with the presence,

absence, or number of the organisms of interest in the sample, or portion thereof.

2-3. (Canceled)

- 4. (Original): The method of claim 1, wherein the detectable molecular probe is not labeled with a detectable moiety.
- 5. (Currently Amended): The method of claim 4, wherein the detectable molecular probe is detected using a detectable antibody that specifically binds to a complex of the detectable molecular probe and the target sequence of the microbial organism of interest probe/target sequence complex.
- 6. (Original): The method of claim 5, wherein the detectable molecular probe is an unlabeled peptide nucleic acid.
- 7. (Original): The method of claim 1, wherein the detectable molecular probe is labeled with a detectable moiety.
- 8. (Original): The method of claim 7, wherein the detectable moiety is selected from the group consisting of: a chromophore, a fluorochrome, a spin label, a radioisotope, an enzyme, a hapten and a chemiluminescent compound.

9-10. (Canceled)

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11. (Currently Amended): The method of claim 1, wherein the solid carrier is selected from the group consisting of: a particle, a bead, a microscope slide, a micro titre plate, and a membrane and an array.

12-14. (Canceled)

1 15. (Original): The method of claim 1, wherein the sample, or portion thereof, is 2 treated with the detectable molecular probe or probes before being contacted with the solid 3 carrier.

| 1 | 16. (Original): The method of claim 1, wherein the sample, or portion thereof, is |
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| 2 | contacted with the solid carrier before being treated with the detectable molecular probe or |
| 3 | probes. |
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| 1 | 17. (Original): The method of claim 1, wherein the sample, or portion thereof, is |
| 2 | simultaneously contacted with both the solid carrier and treated with the detectable molecular |
| 3 | probe or probes. |
| 1 | 18. (Currently Amended): A method for sorting and determining [[an]] a |
| 2 | microbial organism or microbial organisms of interest in a sample or samples; said method |
| 3 | comprising: |
| 4 | treating the sample or samples, or a portion thereof, with one or more detectable or |
| 5 | independently detectable molecular probes wherein the one or more molecular probes are |
| 6 | peptide nucleic acid and are selected such that either: |
| 7 | (i) the detectable probe or probes react with a target sequence of the different |
| 8 | microbial organisms to be determined in a way that produces different |
| 9 | detectable organisms that possess the same stain; or |
| 10 | (ii) the independently detectable probes react with a target sequence of the |
| 11 | different organisms to be determined in a way that produces different |
| 12 | independently detectable organisms that possess an independently |
| 13 | detectable stain; and |
| 14 | contacting the sample or samples, or a portion thereof, with one or more different |
| 15 | types of coded beaded supports, wherein each different type of coded beaded support can |
| 16 | be independently determined in a suitable particle sorter and wherein to the coded beaded |
| 17 | supports have been immobilized one or more antibodies binding partners chosen to select |
| 18 | a particular organism or organisms such that the detectable or independently detectable |
| 19 | organisms become selectively bound to the coded beaded supports as a result of the |

occurrence of specific antibody binding partner interactions;

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sorting the different types of coded beaded supports in a suitable the particle sorter;
and
determining the presence, absence, or number of detectable organisms, or each of

determining the presence, absence, or number of detectable organisms, or each of the independently detectable organisms, immobilized to each different type of coded beaded support and either: (iii) correlating the result with the particular <u>antibody binding partner</u> immobilized to each <u>particle</u> type <u>of coded bead</u> to thereby determine the presence, absence or number of each of the different <u>microbial</u> organisms of interest in the sample, or portion thereof; or (iv) correlating the result with the code for a sample source from which the sample, or portion thereof, was derived to thereby determine the presence, absence or number of each of the <u>microbial</u> different organisms of interest in each different sample, or portion thereof.

19-20. (Canceled)

- 21. (Original): The method of claim 18, wherein the detectable molecular probe is not labeled with a detectable moiety.
- 22. (Currently Amended): The method of claim 21, wherein the detectable molecular probe is detected using an detectable antibody that specifically binds to a <u>complex of the</u> detectable molecular <u>probe and the target sequence of the microbial organism of interest probe/target sequence complex</u>.
- 23. (Original): The method of claim 22, wherein the detectable molecular probe is an unlabeled peptide nucleic acid.
- 24. (Original): The method of claim 18, wherein the detectable molecular probe is labeled with a detectable moiety.
- 25. (Original): The method of claim 24, wherein the detectable moiety is selected from the group consisting of: a chromophore, a fluorochrome, a spin label, a radioisotope, an enzyme, a hapten and a chemiluminescent compound.

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1 26. (Original): The method of claim 18, wherein the independently detectable probes are labeled with independently detectable fluorophores.

27-28. (Canceled)

- 29. (Currently Amended): The method of claim 18, wherein the sample, or portion thereof, is treated with the detectable or independently detectable molecular probe or probes before being contacted with the <u>one or more different types of coded beaded supports solid carrier</u>.
- 30. (Currently Amended): The method of claim 18, wherein the sample, or portion thereof, is contacted with the <u>one or more different types of coded beaded supports solid earrier</u> before being treated with the detectable or independently detectable molecular probe or probes.
- 31. (Currently Amended): The method of claim 18, wherein the sample, or portion thereof, is simultaneously contacted with both the <u>one or more different types of coded beaded supports solid carrier</u> and treated with the detectable or independently detectable molecular probe or probes.

32-34. (Canceled)

35. (Currently Amended): A method for sorting and determining different microbial organisms of interest in a sample; said method comprising:

treating the sample, or a portion thereof, with one or more detectable or independently detectable molecular probes wherein the one or more molecular probes are peptide nucleic acid and are selected such that either:

(i) the detectable probe or probes react with a target sequence of the different

(i) the detectable probe or probes react with a target sequence of the different microbial organisms to be determined in a way that produces different detectable microbial organisms that possess the same stain; or

(ii) the independently detectable probes react with a target sequence of the different microbial organisms to be determined in a way that produces different independently detectable microbial organisms that possess an independently detectable stain; and

contacting the sample, or a portion thereof, with a solid carrier array to which antibodies binding partners have been immobilized at unique identifiable locations such that the detectable or independently detectable microbial organisms are selectively bound to the locations on the array as a result of the occurrence of specific antibody binding partner interactions; and

determining the presence, absence or number of the detectable or independently detectable <u>microbial</u> organisms immobilized at the many different locations of the array and correlating the result with the particular <u>antibody binding partner</u> immobilized to each location on the array to thereby determine the presence, absence or number of the different <u>microbial</u> organisms of interest in the sample.

36-37. (Canceled)

- 38. (Original): The method of claim 35, wherein the detectable molecular probe is not labeled with a detectable moiety.
- 39. (Currently Amended): The method of claim 38, wherein the detectable molecular probe is detected using a detectable antibody that specifically binds to a <u>complex of the</u> detectable molecular <u>proble and the target sequence of the microbial organism of interest probe/target sequence complex</u>.
- 40. (Original): The method of claim 39, wherein the detectable molecular probe is an unlabeled peptide nucleic acid.
- 1 41. (Original): The method of claim 35, wherein the detectable molecular probe 2 is labeled with a detectable moiety.

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| 1 | 42. (Original): The method of claim 41, wherein the detectable moiety is |
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| 2 | selected from the group consisting of: a chromophore, a fluorochrome, a spin label, a |
| 3 | radioisotope, an enzyme, a hapten and a chemiluminescent compound. |

43. (Original): The method of claim 35, wherein the independently detectable probes are labeled with independently detectable fluorophores.

44-45. (Canceled)

- 46. (Original): The method of claim 35, wherein the sample is treated with the detectable or independently detectable molecular probe or probes before being contacted with the solid carrier.
- 47. (Original): The method of claim 35, wherein the sample is contacted with the solid carrier before being treated with the detectable or independently detectable molecular probe or probes.
 - 48. (Original): The method of claim 35, wherein the sample is simultaneously contacted with both the solid carrier and treating with the detectable or independently detectable molecular probe or probes.

49-59. (Canceled)